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Appl. No. 09/524,531
Docket No. 11422/264679

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APPROVED BY	O.G. FIG.	CLASS	SUBCLASS
			DRAFTSMAN

FIG. 1A

```

>muCRAM-1
CAGACATCCCTCGCACATGGCGCTGAGCCGGGCTTGCAGCTTGCTGCTCTTCAGGGCTGCAATGAGGGCATGATAAGTGTGAATTCTCAAAATCCAGCAA
CCGAAACCAGTGGTACATGAAATTGAAAGTGTGAATTGTGCTTGCAGCTTGCTCTTCAGGGCATGATAAGTGTGAATTCTCAAAATCCAGCAA
CAAGTGACCCCTAGGATTGAAATGGAAAGAAATTCCAAGATGCCAACATATGTTGCTTGCAGATGCTGTTGGAAAAACTTCCCTGAGGATCTG
AACAAAGATTCAAGGAGACCTGGCAGGTTGCGACAGATGCTGTTGGAAAGAAATTCCAAGGCAATCTGCTGTGCTCTAGGAGTCTGAAATGACCGAAAAG
GAATGCTGACACGGATCGGATTCAAGGATCGGATTCAGCCATCTAGGCTGTGAGGTGCTGTTCAATGACCGAAAAG
AAGTTGATGAGATTACCATGAGTTAATTGCAAGTGAAGTGAAGGCCAGTGAGCTGCTTCAATGCTGAGGAAATTCTGCTGAGGAAATT
CCAGCCGGCTGTGACTCTGTAGGCCAACACTGGCAAGAACACTGGCACCTGAGGAGACAGGGCACCTCTGGGAGTGTGAGGAAATT
GCCCTCACTACAGCTGGTACCGCAATGATGTTGCCACTGGCACCTGAGGAGACAGGGCACCTCTGGGAGTGTGAGGAAATT
TCCAGAAATTCCATGTAAGCTGGGACTACTGCAATTGCTTCCAGTGGGATTATTGGGATTTGCTGCTTATTG
GACGACTCTGGCAGTACTGCAATTGAAACATTGCTGCTTCCAGTGGGATTTGCTGCTTATTG
GGACATGGAAAGTCTATGATTGAAACATTGCTGCTTCCAGTGGGATTTGCTGCTTATTG
TTCTTGCTGTGATTACGATGGCATTGCTGCTTCCAGTGGGATTTGCTGCTTATTG
CAAGATGGAGAAAGCTATAAGGAGCCCAGGGAGCATGACGGGTGTTAACTACATCCGGACGGAGTGA
GGAGGGGTGACTTCAGACACAATCGTCTTGTATTGCTGACACCTGTGGGAGGACATG
CAAGTACCTCTGGTGAAGCTGGTCAACGGGCTGTGCTTCTGGTACAGGAAACTCTGGTCAAGGGCTTCCAGGAAACTCTGGTCAAGGGCAATT
CGGGCAGAACCTTTTGTITGGGATTAAGGGTGAATGGTCAAGGGCTTCCAGGAAACTCTGGTCAAGGGCAATT
AGCCACAAGAAATAAAAGGAAGCCCTCCTGAAGATGGTCAAGGGCTTCCAGGAAACTCTGGTCAAGGGCAATT
TCATTATGGGATTGGGCTTGGCCAGGGCTGTTAGCTGCCAGGACAACTGGCAGAGGGCTGAGGCGG
TGGACTGTGTTTGGCCAGGGCAACGGGCAACGGCTGATGCCCTGTGACAGTGGCCCCAGGAAGGTTTCAAGG
GGAGGGGTGACTTCAGACACAATCGTCTTGTATTGCTGACACCTGTGGGAGGACATG
CAAGTACCTCTGGTGAAGCTGGTCAACGGGCTGTGCTTCTGGTACAGGAAACTCTGGTCAAGGGCAATT
CGGGCAGAACCTTTTGTITGGGATTAAGGGTGAATGGTCAAGGGCTTCCAGGAAACTCTGGTCAAGGGCAATT
AGCCACAAGAAATAAAAGGAAGCCCTCCTGAAGATGGTCAAGGGCTTCCAGGAAACTCTGGTCAAGGGCAATT
TCATTATGGGATTGGGCTTGGCCAGGGCTGTTAGCTGCCAGGACAACTGGCAGAGGGCTGAGGCGG
TGGACTGTGTTTGGCCAGGGCAACGGGCAACGGCTGATGCCCTGTGACAGTGGCCCCAGGAAGGTTTCAAGG
AGTGGCCTTGCTCCCTGGACCCCTGACCTGGAAATGGTGTGAAATCAGGAACTCTGGTCAAGGGCAATT
CCATCCTGGAGAAATGGTGTGAAATGGTACTGAAGTCTCCCTTCTGATTGGTCTTGTGCTTATTG
CTTGCTGGAAAGGGCTCTGACCCAGGAAACTCTGGTCAAGGGCAATT
ACCTGTCTGTCTTAAGTCTGAAATGGTACTGAAGTGTATTGATTACTGAAAAATTCTGATTATAAAACT
AAAAATTAAACATTCTAAATTGCTAGAGATGTATTGATTATAATGAGTTCAACTTAAGGTAGA
GTAAAATATGCCATACAGTGTTCAAAACGTATTTTATAATGAGTTCAACTTAAGGTAGA
AGGAGGGCTGGCTGCTGAATATTCTCTTGTGTTCAAAAGAAGTTAGCATTTCATAAGAAAACCTAACTCTAAAGAATTC
CTGTCCTGACCAACTGGTGTGCTTAGCTTGTGTTCAAAAGAAGTTAGCATTTCATAAGAAAACCTAACTCTAAAGAATTC

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FIG. 1B

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>mUCRAM-2
CAGAATTCTGGCACGAGGGTCTGGGGCCGACCTACGGTTCTCCCTAAGAGCTAATCTTGCC
GCCACTCGCTTAGGACCTGGACACCGGTCCCCGTCACGGTCAACCCCTTCCACCCCT
CAAAGAACGGACTGCTCAGACACCACCGTCCAGGGCAAGAACCTGGCTGAGACA
CCCCAACGGAGGGACTGACATGGGACAGGAACCCAGGGCTGTGGGCTCAGCTCTG
CTCCGGCGGGGGGAAGTGGCTGGACACCACGGCTCTGGTCTACTACCCAAAGGCTG
CAGGCAGCTGGGGGGGGCTAGATGGCAGGGAGGGCTAGATAAGGCAATGGTTTCTGCA
TGATCTGTGCCCTGGACTATCATATAAGGCAATGGTTTCTGCACTAAAGGCTACA
GTAATAGGTCCAAGAGGGCTATTGGCTTGAAAGGATAAAAGTCAGTGAAGTACTAG
GAAGAAGGTGGACAGGGGGTCTCTGGTCTACTACCCAAAGGCTGAGCTGGAGCTACGGAT
CTGAGATGATAATTCAATTACCAATCAAAATGTTACAGAACGAGTCACTGGGAACTTCTG
GTCAAGGGCTCCGACTGAGCAAGGGCCAGAACCTGGCCACTTCTGTTAGACTGGAA
TGCTGTCTGGCTGAAGTGCCCAGTGGCTTATGACTGGAAAGTGGGAGCTACGGAT
AAGAAGGAAACCGCTCGGGAGTACATCTGGTTAAAGATGGCACAAAGTTGGTAGGGAA
GGCACACACAAACAGCTGAGATTACTGGGAAGCCGGAACTCTGTGACACCCGGCAT
GATGCAAGTGGAGACTTCTGGCTGAAGTGGCTTAAAGGCTACCTTCAAAAGAAACTTCC
TCCAAGTAGATGTTCTAACATAAGCGGCATCATAGGAACGGGTGTTAGGCAACACAA
TGTGCTCTGGCACATGGCTAGCTCAGGGAAAGGCTACCTTCAAAAGAAACTTCC
TCCTGCATCTAAAGTCACTACGGATGGCCAAACCAGTTGTCACTAGTTAGGCAACACAA
ATTCCAGTTGGCTGCCAAACCAGTTGTCACTAGTTAGGCAACACAAATCTGGGAA
CTTGGAAAGTGAAGAAATATGAAAGGAGTTCACTAGAACTTCAAAAGGCTACCTA
ATTITAAGCAAAACGTCTTGTGCACTAACTGCAAGTGGGAAATCTGTTGAGTTGACT
TGTTAATATTCTGGTTTGTGAGTTGACTATACACTGACTGAGTGGAAATTTAAAGA
CTTGAATCTGGCACAGAAATGAAAGGAAATATGAAAGGAAATATGAAAGGAAATTT
CTTAACTCCAGAACGAAATATAAAAGGAAATATAAAAGGAAATATAAAAGGAAAT

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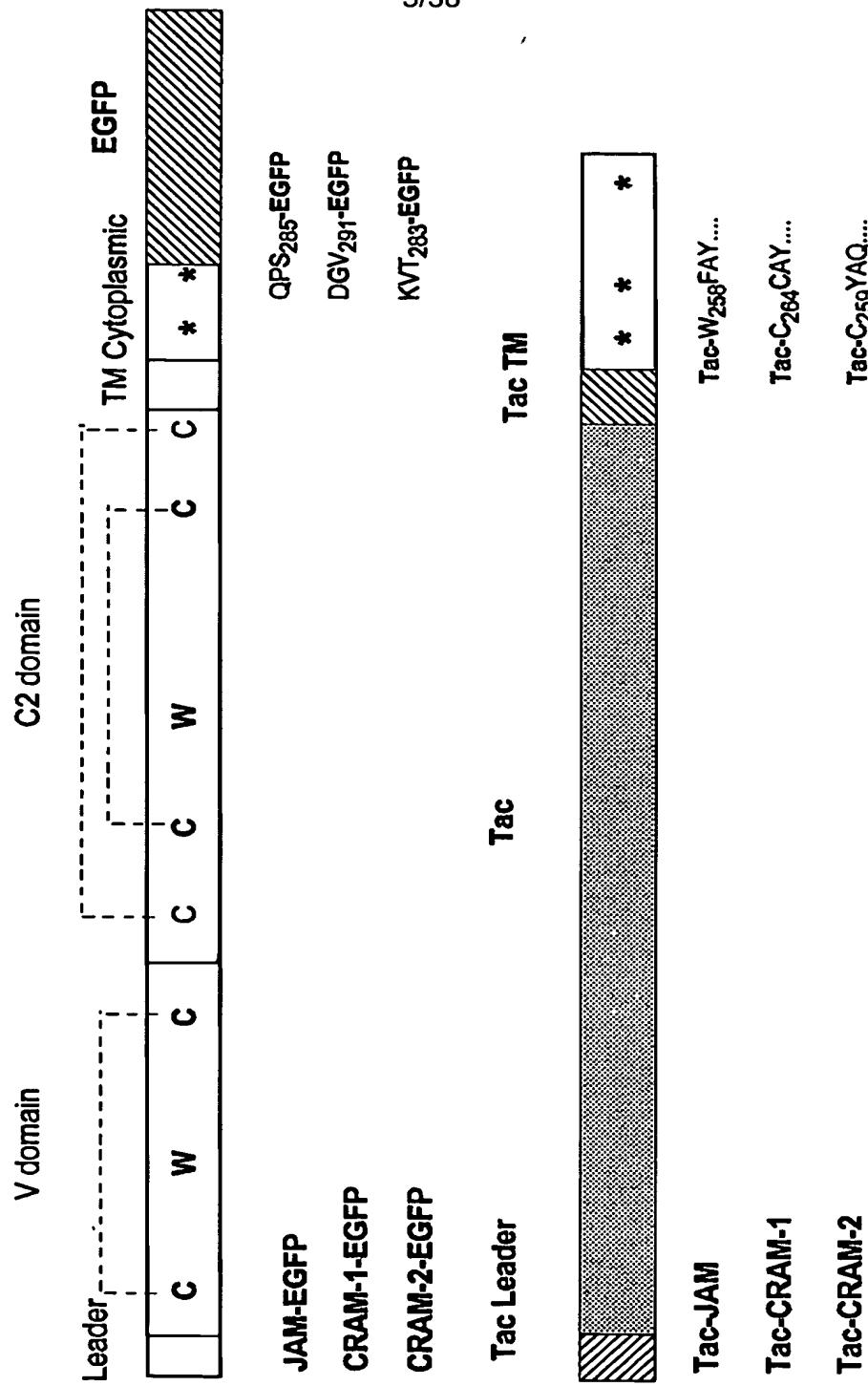


FIG. 2A



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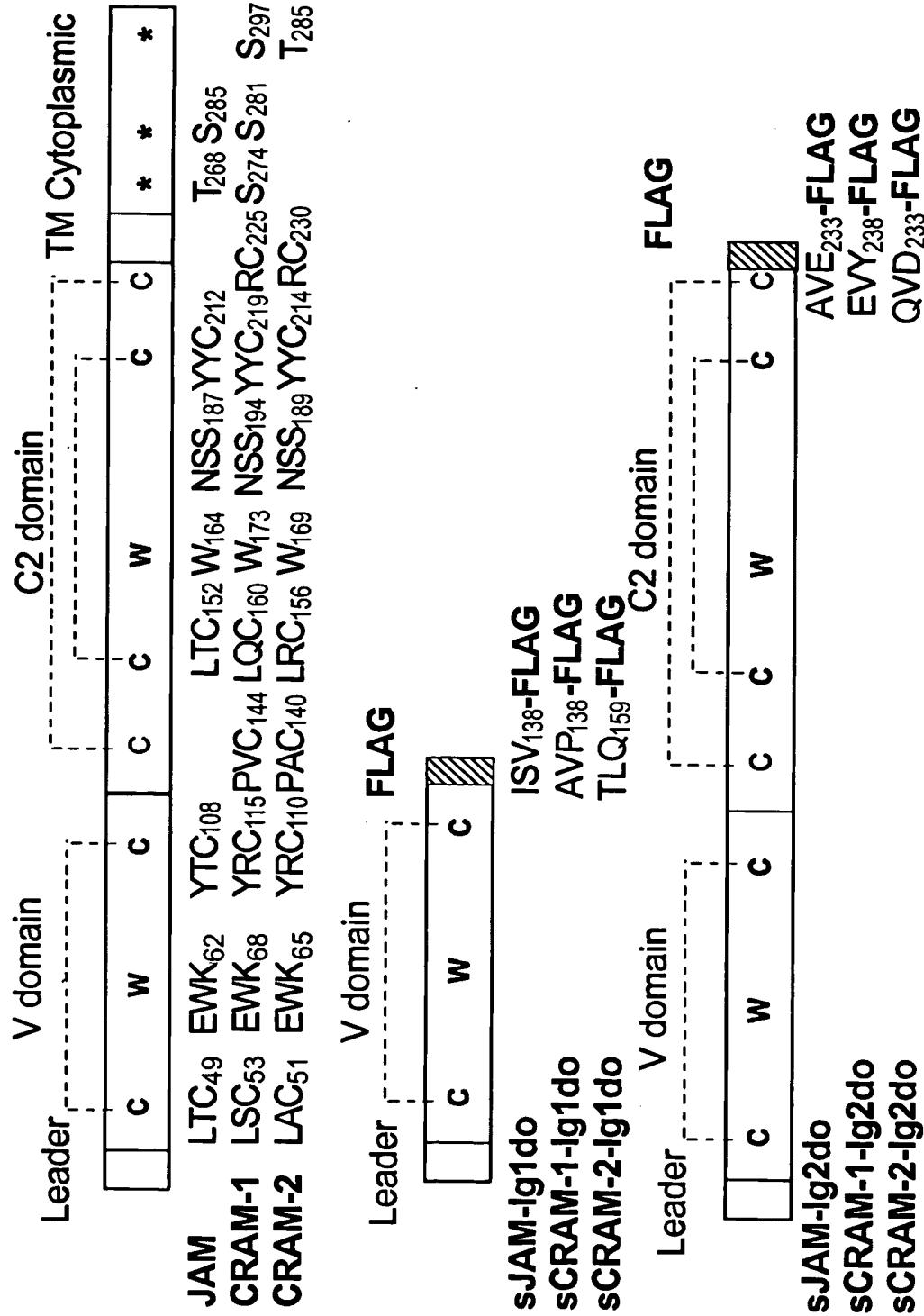


FIG. 2B

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MALSRRIRLR LYARILPHFFL LILFRGCMIE AVNLKSSNRM PVVHEFESVE
MARSPOGL IMLLLHHLI VALDYHKANG FSASKDHROE VTVTIEFQEAI

LSCIIITHSQT SDPRIEWKKI QDGQTTYVVF DNKIQGDLAG RTDVFGKTSL
LACK-TPKKT TSSRLEWKVV GQG-VSLVYY QQALQGDFKD RAEMID-FNI

RIWNVTRSDS ATYRCEVVAL NDR-KEVDEI TIELIIVQVKP VTPVCRIPAA
RIKNVTRSDA GEYRCEVSAP TEQGQNLQED KVMLEVVLVAP AVPACEVPTS

VPVVGKTATLQ CQESEGYPRP HYSWYRNDVP LPT-DSRANP RFQNSSSHVN
VMTGSVVELR CQDKEGNPAP EYIWFKDGTS LLG-NPKGGT H-NNSSY'TNE

SETGTLVFNA VHKKDDSGQYY CIASNDAAGAA RCEGQDMEVY DLNIAGITIG
HESGILQFNM ISKMDSGEYY CEARNSVGHR RCPGKRMQVD VLNIISGIAT

VLVVLIVLAV ITMGICCCAYR RGCFISSLQD GESYKSPGKH DGVNYIRTSE
VVVAFVISV CGLGTCYAQR KGKF--SKET SFQKGSPASK ---- VTMG

EGDFRHKSFF VI
ENDFRHTKSF II

FIG. 3

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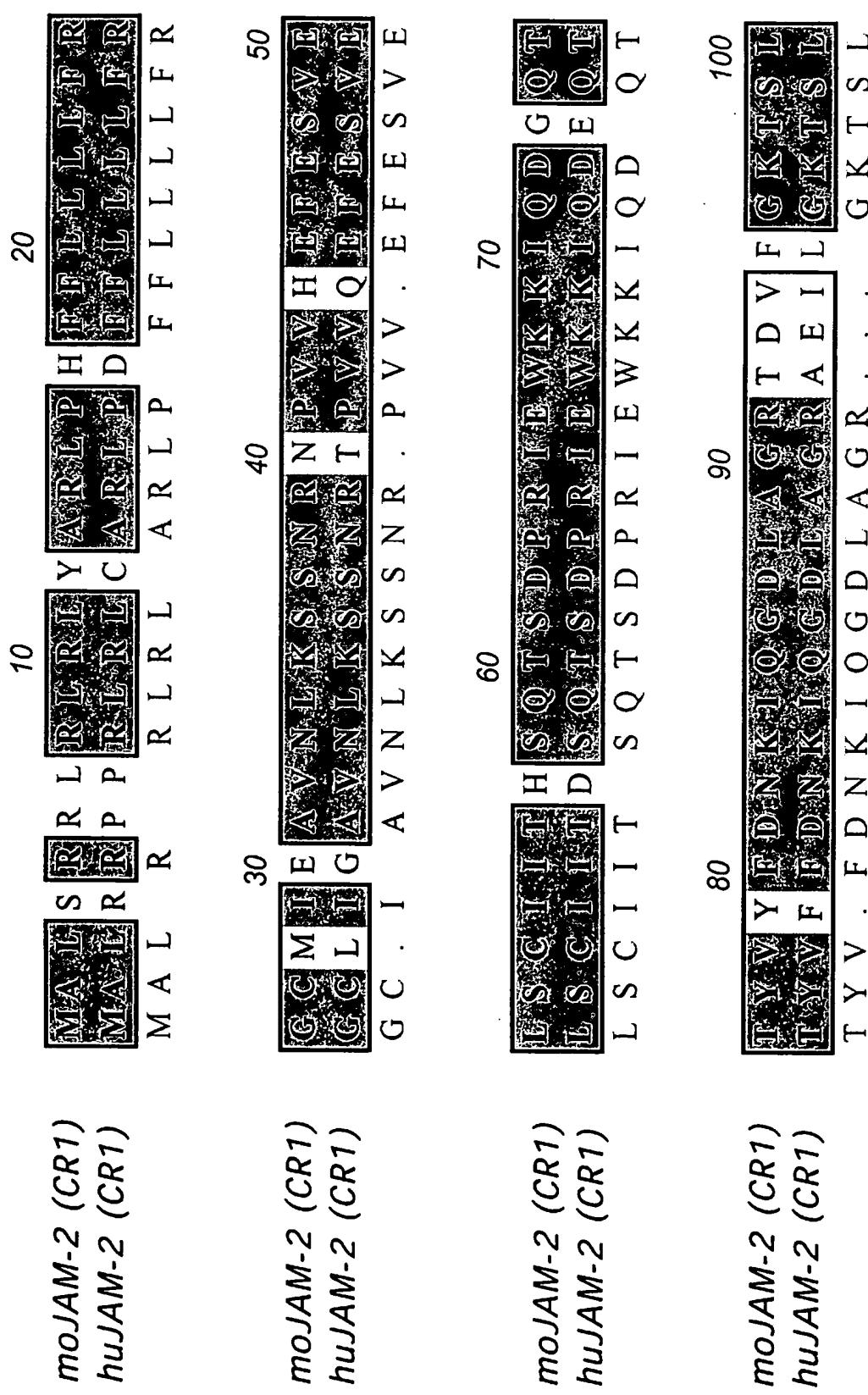
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FIG. 4A

Murine and human JAM-2 (CRAM-1)





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FIG. 4B

Murine and human JAM-2 (CRAM-1)

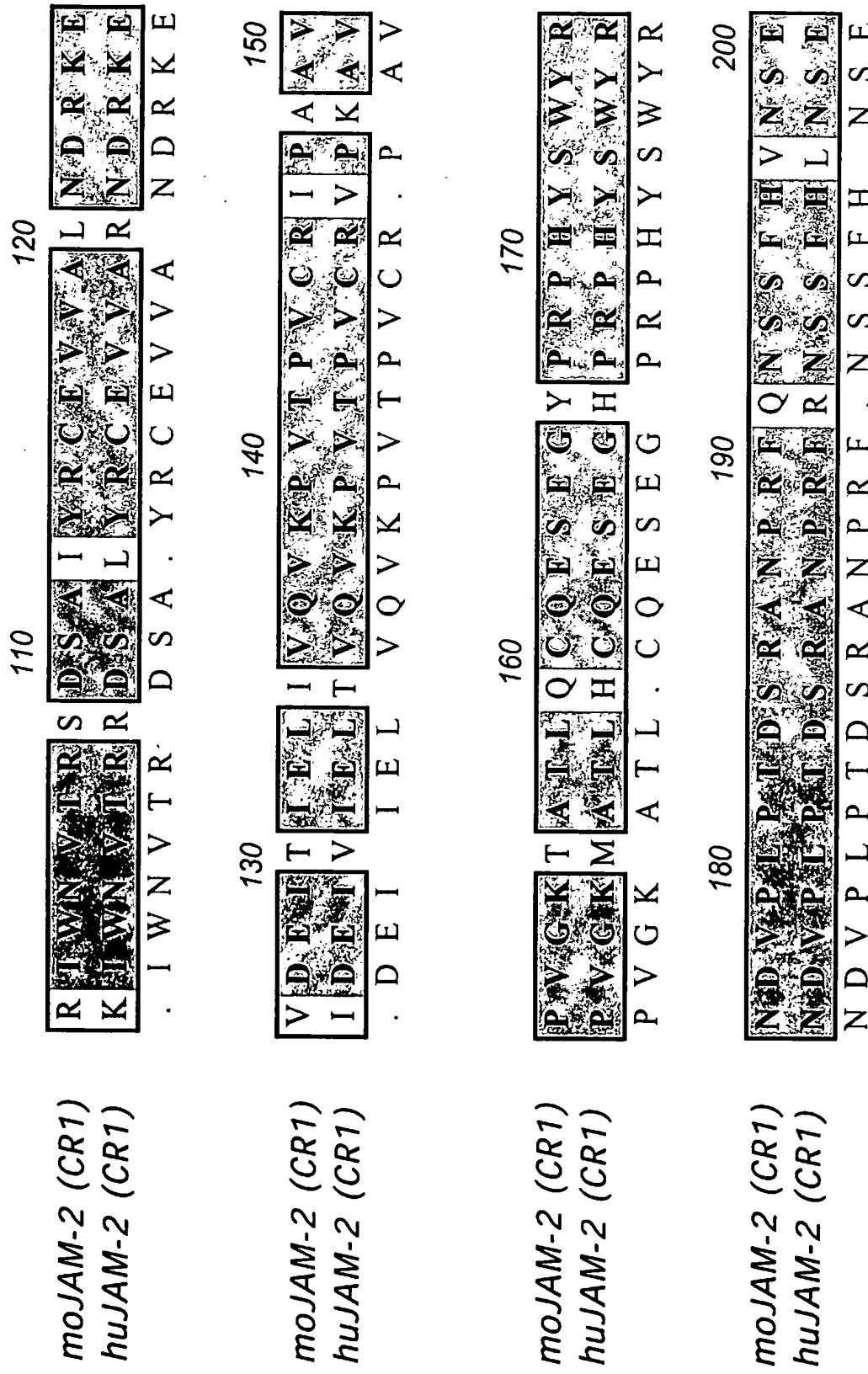


FIG. 4C

Murine and human JAM-2 (CRAM-1)

210

<i>moJAM-2 (CR1)</i>	N	A	V	H	K	D	D	S	G	Q	V	A	C	T	A	S	N	D	A				
<i>huJAM-2 (CR1)</i>	I	G	T	V	E	T	A	V	H	K	D	D	S	G	O	Y	C	I	A	S	N	D	A

T G T L V F . A V H K D D S G O Y Y C I A S N D A

220

230

<i>moJAM-2 (CR1)</i>	G	A	A	R	G	Q	D	M	E	V	Y	D	E	N	T	A	G	I	I	G	G	V	L
<i>huJAM-2 (CR1)</i>	G	S	A	R	C	E	Q	E	M	E	N	D	E	N	L	G	G	I	J	G	G	V	L

G . A R C E Q . M E V Y D L N I G I I G G G V L

240

<i>moJAM-2 (CR1)</i>	V	V	L	A	V	I	T	M	G	I	C	C	A	Y	R	R	G	C	F	I	S	S
<i>huJAM-2 (CR1)</i>	V	V	L	A	V	I	T	L	G	I	C	C	A	Y	R	R	G	Y	F	I	N	N

V V L V L A . I T . G I I C C A Y R R G F I

250

<i>moJAM-2 (CR1)</i>	260	V	V	L	A	V	I	T	M	G	I	C	C	A	Y	R	R	G	C	F	I	S
<i>huJAM-2 (CR1)</i>	270	V	V	L	A	V	I	T	L	G	I	C	C	A	Y	R	R	G	Y	F	I	N

V V L V L A . I T . G I I C C A Y R R G F I

260

<i>moJAM-2 (CR1)</i>	280	K	Q	D	G	E	S	Y	K	S	P	G	K	H	D	G	V	N	Y	I	R	T	S
<i>huJAM-2 (CR1)</i>	290	K	Q	D	G	B	S	Y	K	N	P	G	K	P	D	G	V	N	Y	I	R	T	D

K Q D G E S Y K P G K D G V N Y I R T E E G

280

290

300

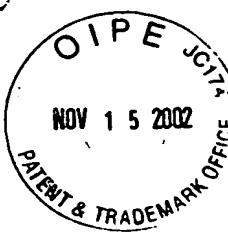
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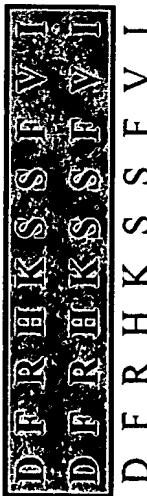
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FIG. 4D Murine and human JAM-2 (CRAM-1)

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310



320

350

340

330

moJAM-2 (CR1)
huJAM-2 (CR1)

360

370

moJAM-2 (CR1)
huJAM-2 (CR1)

400

390

380

moJAM-2 (CR1)
huJAM-2 (CR1)



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FIG. 5A

Murine and human CRAM-2 (JAM-3)

10 20

huJAM-3(CRAM-2) M A R S P Q G L L M L L H Y L I V A L D Y H K

30 40 50

huJAM-3(CRAM-2) A N G F S A S K D H R Q E V T V I E F Q E A I L A

60 70

huJAM-3(CRAM-2) C K T P K T T S S R L E W K K V G Q G V S L V Y

80 90 100

huJAM-3(CRAM-2) Y Q Q A L Q G D F K D R A E M I D F N I R I K N V

moJAM-3(CRAM-2) Y Q Q A L Q G D F K D R A E M I D F N I R I K N V

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FIG. 5B

Murine and human CRAM-2 (JAM-3)

huJAM-3(CRAM-2) moJAM-3(CRAM-2)	110 	120 	
huJAM-3(CRAM-2) moJAM-3(CRAM-2)	130 	140 	150
huJAM-3(CRAM-2) moJAM-3(CRAM-2)	160 	170 	180
huJAM-3(CRAM-2) moJAM-3(CRAM-2)	190 	200 	

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FIG. 5C

Murine and human CRAM-2 (JAM-3)

<i>huJAM-3(CRAM-2)</i>	T V S K L D T G E Y S C F A R N S V G Y R R	220
<i>moJAM-3(CRAM-2)</i>	S R N M I S K M D S G E Y Y C E A R N S V G H R R	
Q F N . S K . D . G E Y C E A R N S V G R R		
210		
<i>huJAM-3(CRAM-2)</i>	C P G K R M Q V D D T N E S G I I A A V V V V A L	250
<i>moJAM-3(CRAM-2)</i>	C P G K R M Q V D V I N I S G I I A T V V V V A F	
C P G K R M Q V D L N I S G I I A . V V V V A		
230		
<i>huJAM-3(CRAM-2)</i>	V I S V C G L G C Y A Q R K G Y F S K E T S F Q	270
<i>moJAM-3(CRAM-2)</i>	V I S V C G L G V C Y A Q R K G Y F S K E T S F Q	
V I S V C G L G C Y A Q R K G Y F S K E T S F Q		
260		
<i>huJAM-3(CRAM-2)</i>	K S N S S S K A H T M S E N D F K H T K S H I I I	300
<i>moJAM-3(CRAM-2)</i>	K G S P A S K V H T M G E N D F R H T K S H I I I	
K . S K T T M E N D F . H T K S F I I I		
280		
<i>huJAM-3(CRAM-2)</i>		290
<i>moJAM-3(CRAM-2)</i>		

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>Nucleic acid sequence huCRAM-1/ huJAM-2
ATGGCGCTGAGGCCACCGCAGTCCGGCTCTGCCTCGGC
TGCCTGACTTCTTCCTGCTGCTTTCAAGGGCTGCCTGAT
AGGGGCTGTAAATCTCAAATCCAGCAATCGAACCCCAGTGGTA
CAGGAATTGAAAGTGTGGAACTGTCTGCATCATTACGGATT
CGCAGACAAGTGACCCAGGATCGAGTGGAAAGAAAATTCAAGA
TGAACAAACACATATGTGTTTGACAACAAAATTCAAGGGA
GACTTGGCGGGTCTGAGAAATACTGGGAAAGACATCCCTGA
AGATCTGGAATGTGACACGGAGAGACTCAGCCCTTATCGCTG
TGAGGTCGTTGCTCGAAATGACCGCAAGGAAATTGATGAGATT
GTGATCGAGTTAACTGTGCAAGTGAAGCCAGTGACCCCTGTCT
GTAGAGTGCCGAAGGCTGTACCACTAGGCAAGATGGCAACACT
GCACTGCCAGGAGAGTGAGGGCCACCCCCGGCCTCACTACAGC
TGGTATCGCAATGATGTACCACTGCCACGGATTCCAGAGCCA
ATCCCAGATTGCAATTCTTCTTCCACTTAAACTCTGAAAC
AGGCACTTGGTGTTCAGTGTGTTCAAAAGGACGACTCTGGG
CAGTACTACTGCATTGCTTCAATGACGCAAGGCTCAGCCAGGT
GTGAGGAGCAGGAGATGGAAGTCTATGACCTGAAACATTGGCGG
AATTATTGGGGGGTTCTGGTTGTCTGACTGGCCCTG
ATCACGTTGGGCATCTGCTGTGCATAACAGACGTGGCTACTTCA
TCAACAATAAACAGGATGGAGAAAGTTACAAGAACCCAGGGAA
ACCAGATGGAGTTAACTACATCCGCACTGACGAGGAGGGCGAC
TTCAGACACAAGTCATCGTTGTGATCTGAGACCCGGGTGTGG
CTGAGAGCGCACAGAGCCGACGTGCACATACCTCTGCTAGAA
ACTCCTGTCAAGGCAGCGAGAGCTGATGCACTCGACAGAGCTA
GACACTCTTCAAAGCTTCTGTTGGCAAGGTGACCACTACTC
TTTACTCTACAAGCCCAGGAAAGAGAAATTCTCAAGAGG
ACCCGGAAATATAACCCCAAGGAACCAAACGGTGCCTCAC
TGAGGTTGGGTCTTAATTGTTTGGCCTGATTCCCATGAA
AATAAGGGGTCTTAAGAGTTGGTACGTAAAACCCCCCGCTT
GGGCCTGGAAACCACATGTTACCACCTGCGTAAAAAAA
AAAAAA

FIG. 6A



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>huCRAM-1/huJAM-2 complete

MALRRPPRLRLCARLPDFFLLLLFRGCLIGAVNLKSSNRTPVV
QEFESVELSCIITDSQTSDPRIEWKKIQDEQTTYVFFDNKIQG
DLAGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEIDEI
VIELTVQVKPVTPVCRVPKAVPGKMATLHCQESEGHPRPHYS
WYRNDVPLPTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSG
QYYCIASNDAGSARCEEQEMEVYDLNIGGIIGGVLVVLAVLAL
ITLGICCCAYRRGYFINNKQDGESYKNPGKPDGVNYIRTDEEGD
FRHKSSFVI

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>huCRAM-2/huJAM-3 partial

RAEMIDFNIRIKNVTRSDAGKYRCEVSAPAEQGQNLEEDTVTL
EVLVAPAVPSCEVPSSALSGTVVELRCQDKEGNPAPEYTWFKD
GIRLLENPRILGSQSTNSSYTMNTKTGTLOFNTVSKLDTGEYSC
EARNSVGYRRCPGKRMQVDDLNISGIIAAVVVVALVISVCGLG
VCYAQRKGYFSKETSFQKSNSSSKATTMSENFKHTKSFI

FIG. 6B

FIG. 7A

Y R C X A S1
TAY AGN TGY NNN GCY TCY AA

Y Q C X A S2
TAY CRG TGY NNN GCY AGY AA

Y Q C X A S1
TAY CRG TGY NNN GCY TCY AA

Y Y C X A S2
TAY TAY TGY NNN GCY AGY AA

Y Y C X A S2
TAY TAY TGY NNN GCY TCY AA

FIG. 7B

t-end/B16
t-end

B16
t-end/B16



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gacattccctcgacatggcgctggccggcttcgacttgcgttatcgactgtacgcggctg 60
M A L S R R L R L Y A R L 15
 cttcaactttcttcgtgtttcagggttcgtatgtatggcattgtaaatctcaa 120
P H F F L L F R G C M I E A V N L K 35
 tccagcaaccggatggtaatcatgtatggaaatgtggatttgtttgttatcat 180
S S N P V V H E F E S V E L S (C) I I 55
 acgactcacagacaatggacccttaggatgtatggaaatccaagaatggccaaacc 240
T H S Q T S D P R I E W K K I Q D G Q T 75
 acatatgttatatttgcacaacaaggattcaaggaggacctggcaggatgtttt 300
T Y V Y F D N K I Q G D L A G R T D V F 95
 ggaaaaactcccttggatctggatgtgacacgatcggttatcgccatctatcgctgt 360
G K T S L R I W N V T R S D S A I Y R (C) 115
 gaggtcggttcgtatggaccggaaaatggatggattaccatgtggatatttgtg 420
E V V A L N D R K E V D E I T I E L I V 135
 caagtgaaggccaggatggaccctgttcgtatccaggccgttatgttagggaaagacg 480
Q V K P V T P V C R I P A A V P V G K T 155

FIG. 8A



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gcaacactgcaggccaaaggaggcgagggttatccccggctcaactacaggctggtaccgc 540
A T L Q (C) Q E S E G Y P R P H Y S W Y R 175
aatgatgtgccactcctacaggattccaggccaaatcccaggttccagaattccctctttc 600
N D V P L P T D S R A N P R F Q N S S F 195
catgtgaactcggagacaggcactctggttttcaatgtctgtccacaaggacgactctggg 660
H V N S E T G T L V F N A V H K D D S G 215
cagtaactgcattgcattccaaatgcggcagggtcaggccagggttgtagggcaggacatg 720
Q Y Y (C) I A S N D A G A A R C E G Q D M 235
gaagtctatgattgaacattgtctggattattgtgggatattgtgggagggtccattttttt 780
E V Y D L N I A G I I G V L V V L I V 255
cttgctgtgattacgatggccatctgtgcgtacagacgaggctgtgttcatcaggcagt 840
L A V I T M G I C C A Y R R G C F I S S 275
aaacaagatggagaaaaggctataaggaggccaggatgacgggttaactacatccgg 900
K Q D G E S Y K S P G K H D G V N Y I R 295
acgagtgtggagggtgacttcagacacaaatcgttccattgtggct 960
T S E E G D F R H K S S F V I * 310

FIG. 8B



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FIG. 8C



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APPROVED	O.G. FIG.	SUBCLASS
BY	CLASS	
CRAFTSMAN		

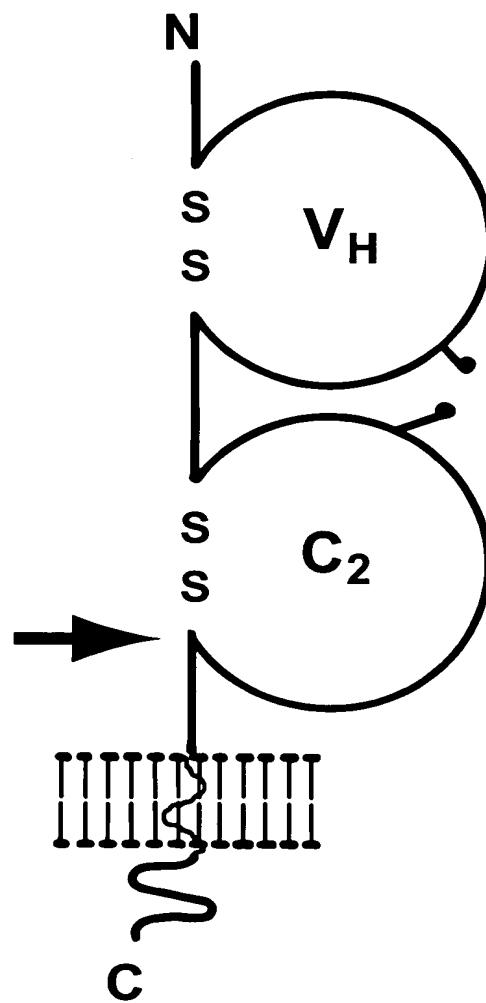


FIG. 8D



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FIG. 9A

APPROVED BY	O.G. FIG. CLASS	SUBCLASS
DRAFTSMAN		

20

MAPSPRIYRILPHEELFR
MARSPOGLMLLHLVALDY
MGTGKAGRKLLLEFTSMILG
JAM Y R C 1 L h f

40

GCMIEAVNLSKSNRNPVVHEFESVE
HKANGFSASKDHROEVTVIEEFEQEAII
SLWQGKGSMYTAAQSDVQVPEMESIK
L W Q G K G S M Y T A Q S D V Q V P E M E S I K
* L S C I I T H S Q T S D P R I E W K K I Q D G Q E

60

LACK-TPKKTTSRLEWKKVGG-V
LTCT--TYSGFSSPRVEWKFVQGSTT
LC T S T S S P R E W K K Q C T
* L S C I I T H S Q T S D P R I E W K K I Q D G Q E

100

TYVIVYEDNKIQGDLAGRTDMEVGKTSI : 100
TYVIVYQQALOGDFKDRAEMID-FNI : 95
ALVCYNSQITAPYADRVSSEGI : 93
IVM add

CRAM - 1 ::
CRAM - 2 ::
JAM ::
Consensus ::

CRAM - 1 ::
CRAM - 2 ::
JAM ::
Consensus ::

CRAM - 1 ::
CRAM - 2 ::
JAM ::
Consensus ::

CRAM - 1 ::
CRAM - 2 ::
JAM ::
Consensus ::



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FIG. 9B

APPROVED	O.G. FIG.
BY	CLASS
DRAFTSMAN	SUBCLASS

* 120

R I W N V T R S D S A I Y R C E V V A L N D R - K
R I K N V T R S D A G E Y R C E V S A P T E Q Q Q
T F S S V T R K D N G E Y T C M V S - - E E G G Q
r i n V T R S D q e y r c e v s a q q

*

E V D E I T I E L I V O V K P V T P V C R I P A A
E N I Q E D K V M L E V L V A P A V P A C E V P T S
N Y G E V S I H L T V L V P P S K P T I S V P S S
E L V I V P C P C P

140

*

V P M G K T A T L Q C Q E S E G Y P R P H Y S W Y
V M T G S V V E L R C D K E G N P A P E Y I W F
V T H G N R A V L T C S E H D G S P P S E Y S W E
V G a L C a L G

160

*

R N D Y P L P T - D S R A N P R E Q N S S F H V N : 198
K D G T S L I G - N P K G G T H - N N S S Y T N E : 193
K D G E S M I T A D A K K T R A E H N S S E T E : 191
k g s 1 t k f n s s t

200

CRAM - 1 ::
CRAM - 2 ::
JAM ::
Consensus ::

CRAM - 1 ::
CRAM - 2 ::
JAM ::
Consensus ::

CRAM - 1 ::
CRAM - 2 ::
JAM ::
Consensus ::

CRAM - 1 ::
CRAM - 2 ::
JAM ::
Consensus ::



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APPROVED BY	O.G. FIG. CLASS	SUBCLASS
		DRAFTSMAN

FIG. 9C

220*

CRAM - 1 CRAM - 2 JAM Consensus

240

+

CRAM - 1 CRAM - 2 JAM Consensus

DAGAA-	RCEGQDM	EVYDLDNIAGIIG
SVGHR-	RCPGKRM	QVDVLNISGIIA
GYGTAMR	SSEAAHMDA	VELNVGGIVAGI

260

CRAM - 1 CRAM - 2 JAM Consensus

300

CRAM-1 CRAM-2 JAM Consensus

ISSKQDGESYKSPGKHDGVNYTRTS	:	297
--SKETSFQKGSPASK-----VTTM	:	285
--ETTKKGTAAPGKKVIYSQPSSTR	:	287
SPqkk		T
sk		t



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APPROVED	O. G. FIG.	CLASS	SUBCLASS
BY			
DRAFTSMAN			

FIG. 9D

CRAM - 1
CRAM - 2
JAM
Consensus

E	G	D	F	R	H	K	S	S	F	V	I
E	G	D	F	R	H	T	K	S	F	I	I
E	G	N	D	E	R	H	T	K	S	F	L
E	G	E	N	D	E	R	H	T	K	S	V
E	G	E	N	D	E	R	H	T	K	S	F



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APPROVED	O.G. FIG.	
	CLASS	SUBCLASS
BY	DRAFTSMAN	

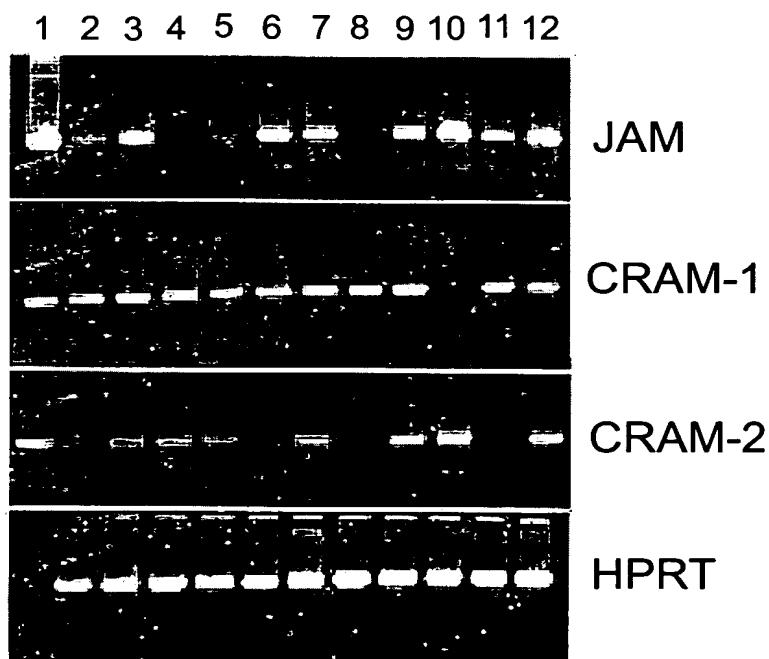


FIG. 10A

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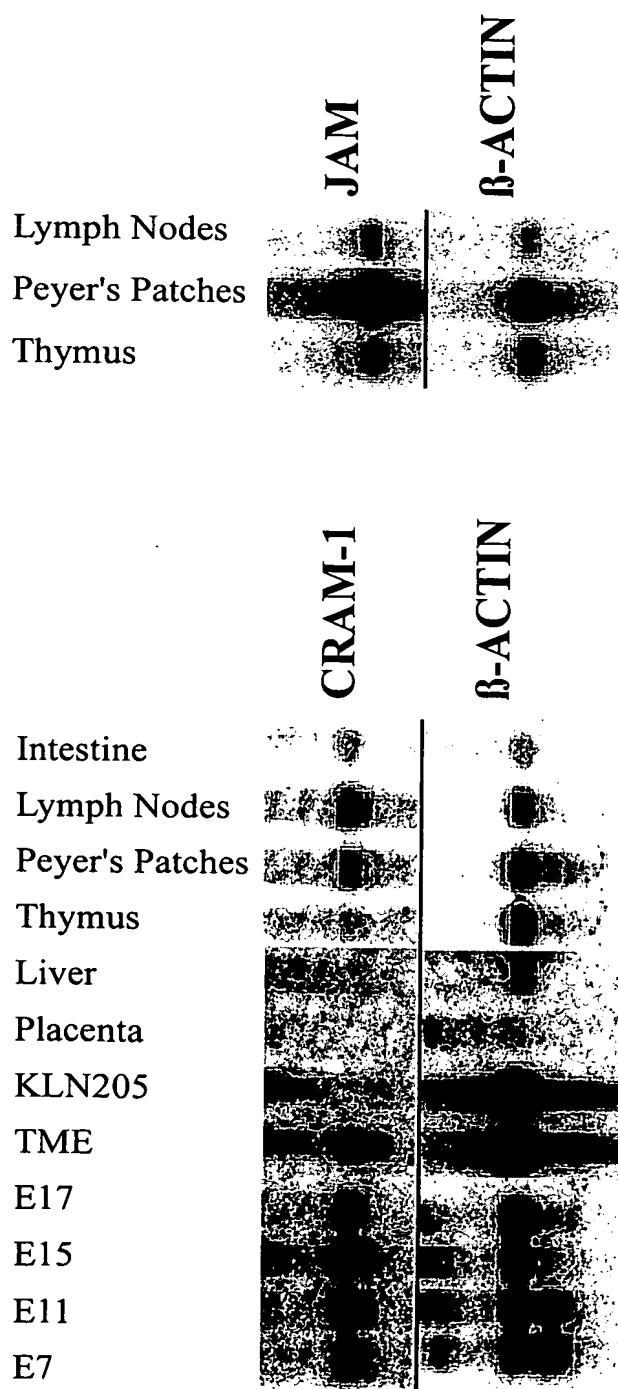
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APPROVED	O.G. FIG. 3.	SUBCLASS
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FIG. 10B



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APPROVED	O.G. FIG.
BY	CLASS
CRAFTSMAN	SUBCLASS

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FIG. 11A

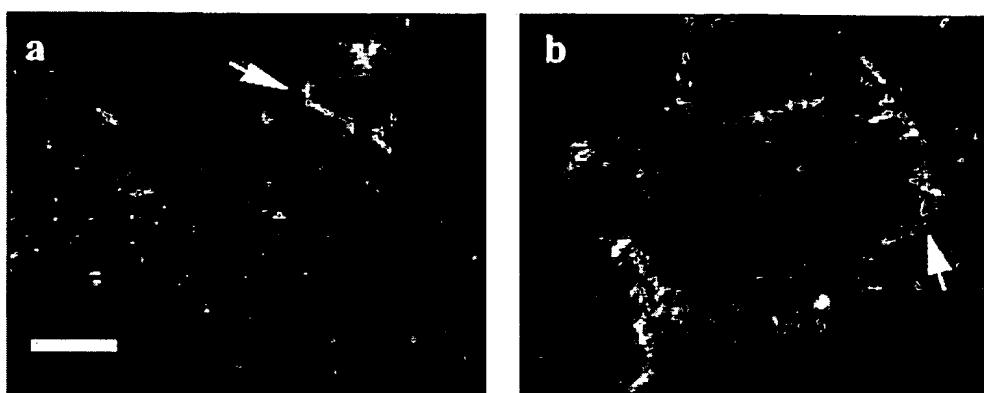
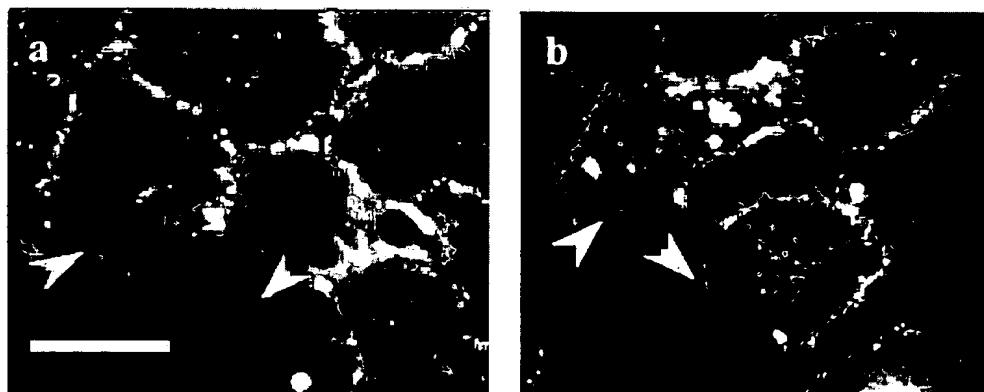


FIG. 11B



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APPROVED	O.G. FIG.	SUBCLASS
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DRAFTSMAN		

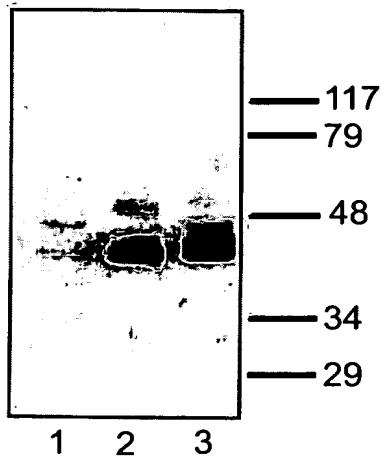


FIG. 11C

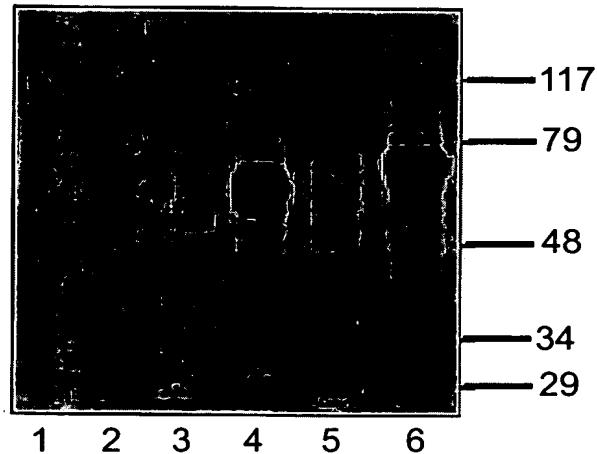


FIG. 11D

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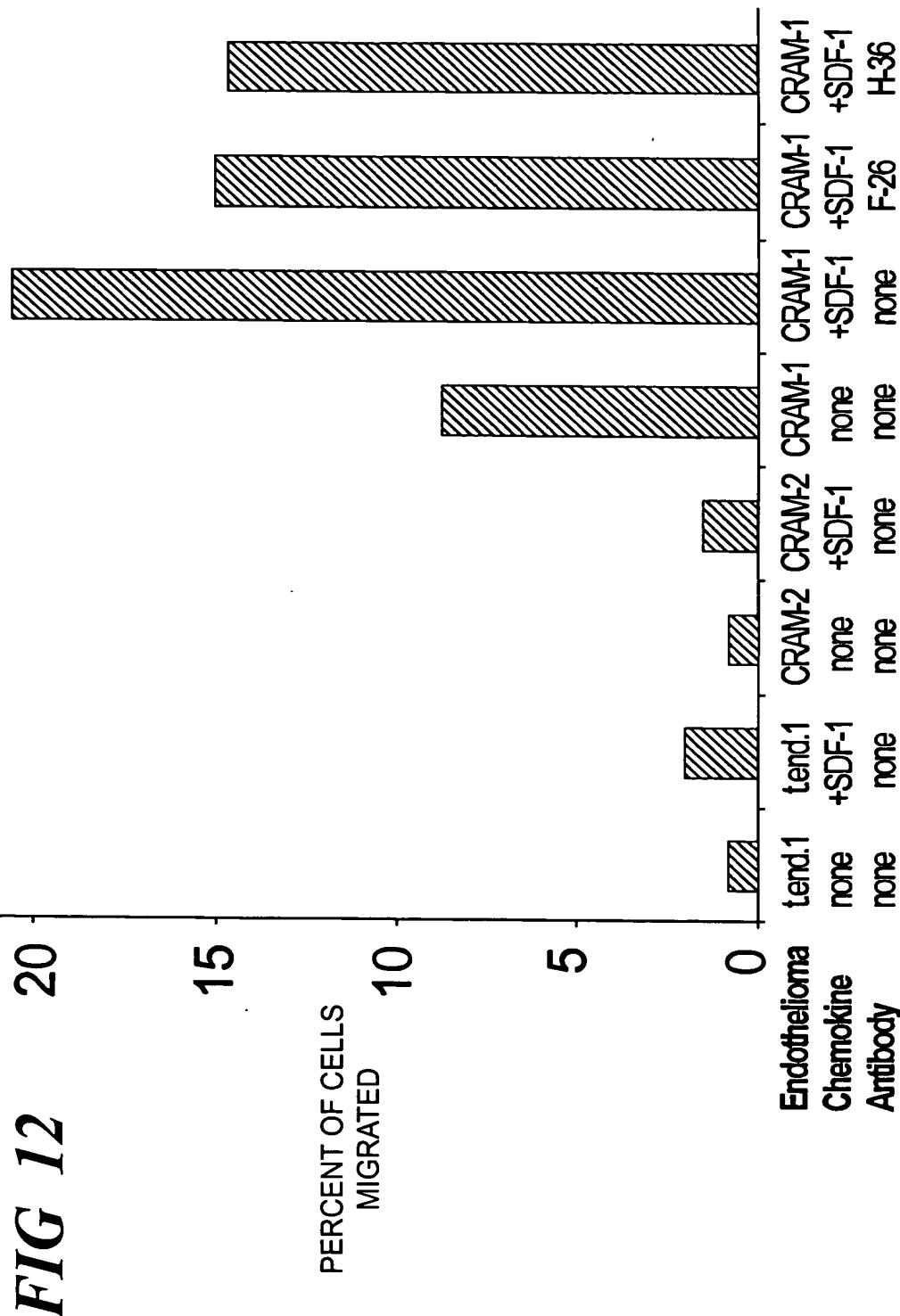
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APPROVED	O.G. FIG.
BY	CLASS
DRAFTSMAN	SUBCLASS

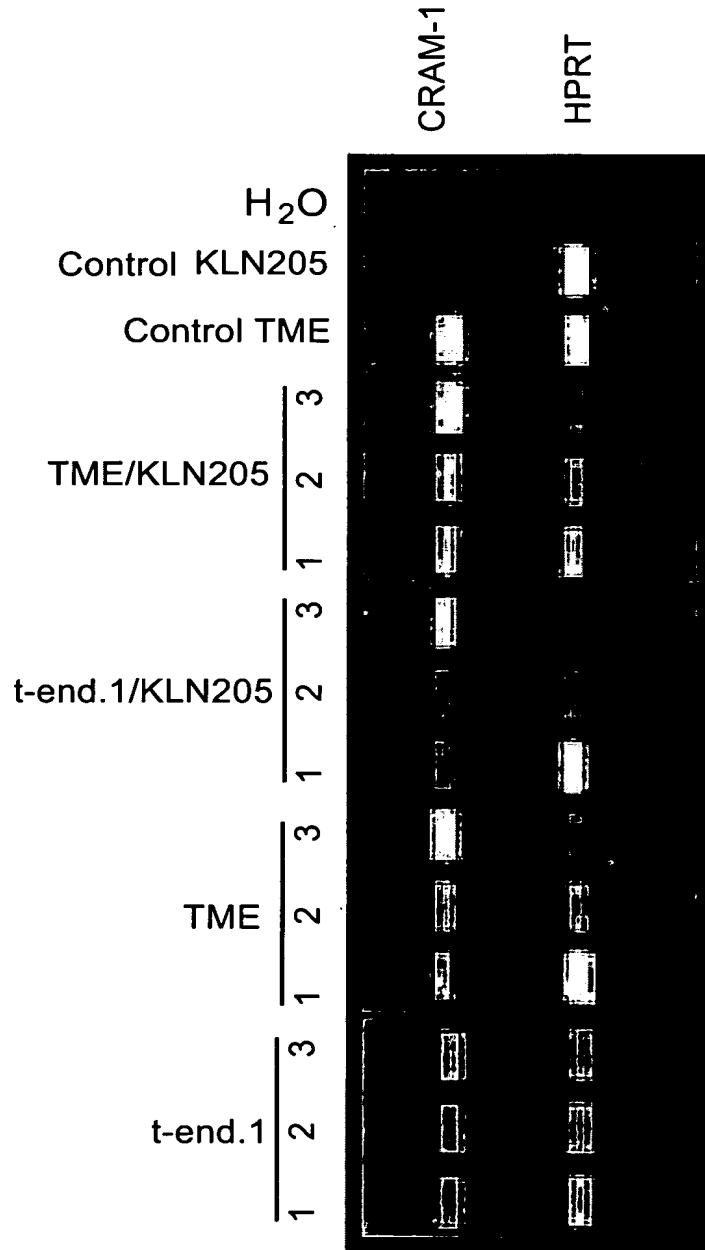


FIG. 13



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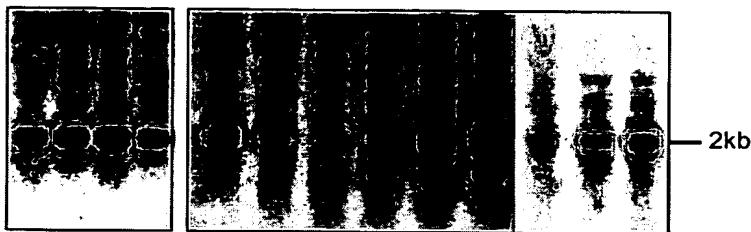
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APPROVED	O.G. FIG.
BY	CLASS
DRAFTSMAN	SUBCLASS

FIG. 14

a



b



c



Day 7 pc
Day 11 pc
Day 16 pc
Day 17 pc

Heart
Lung
Liver
Spleen
Kidney
Testis
Thymus
Peyer's patches
Lymph node



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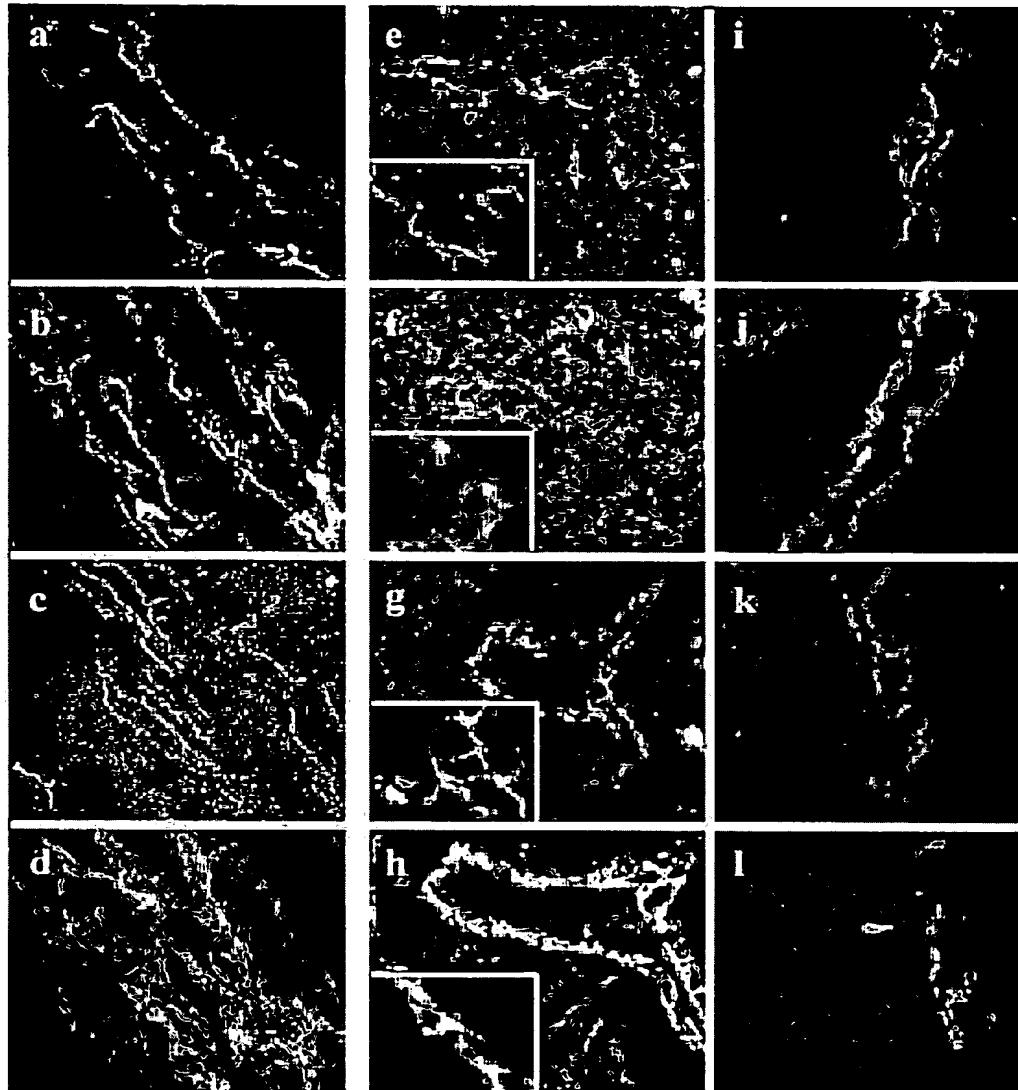


FIG. 15

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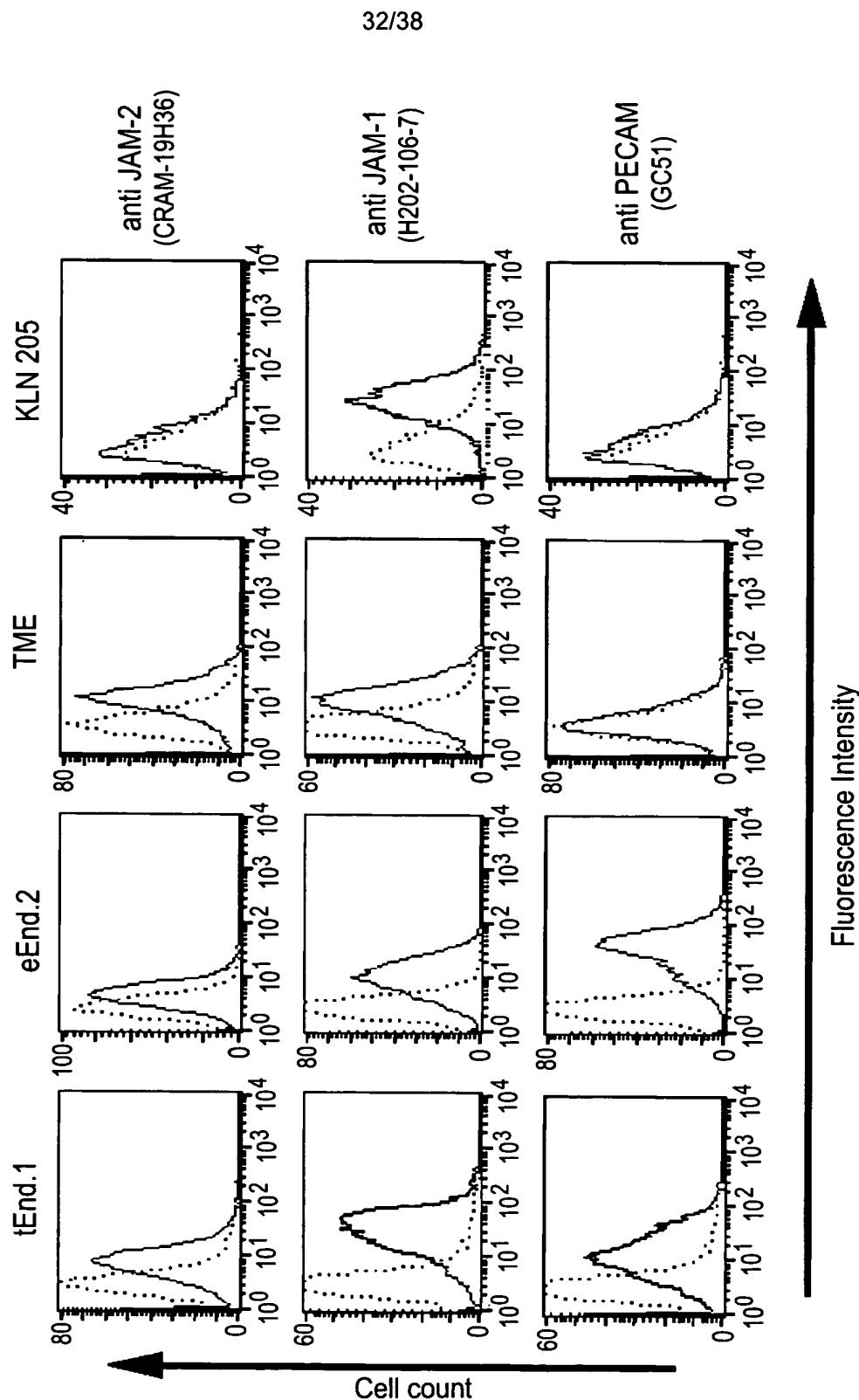
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DRAFTSMAN	SUBCLASS

FIG. 16A



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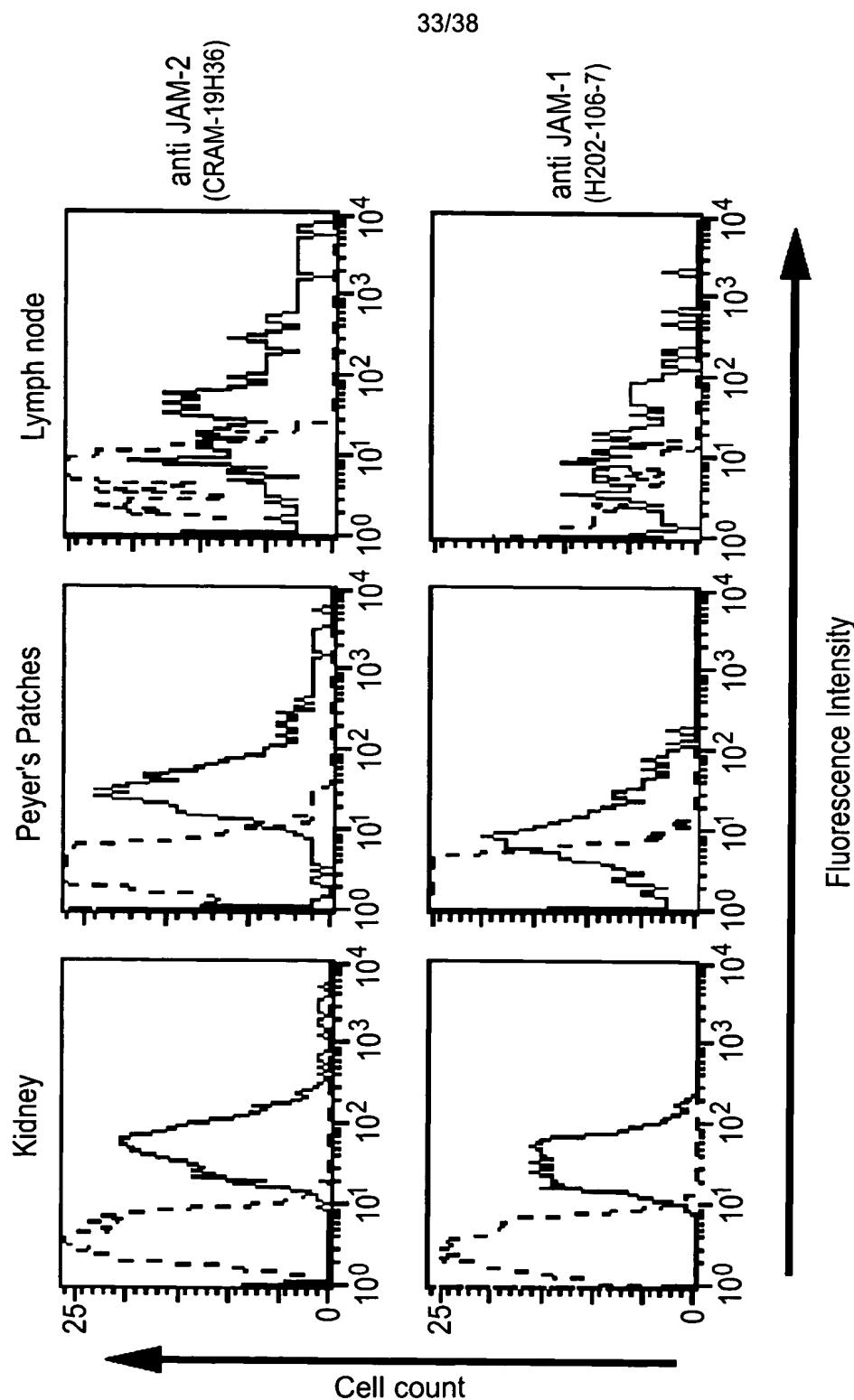
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FIG. 16B





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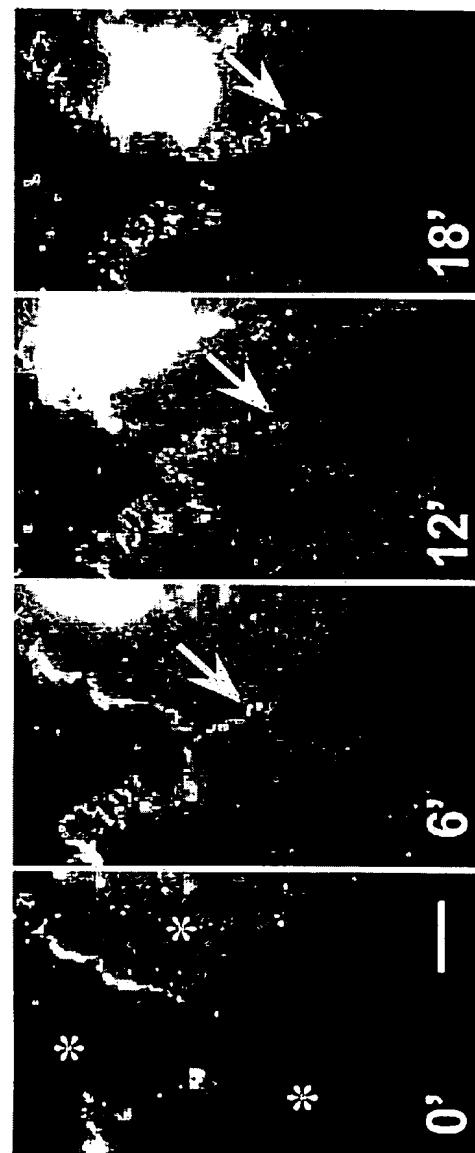


FIG. 17A

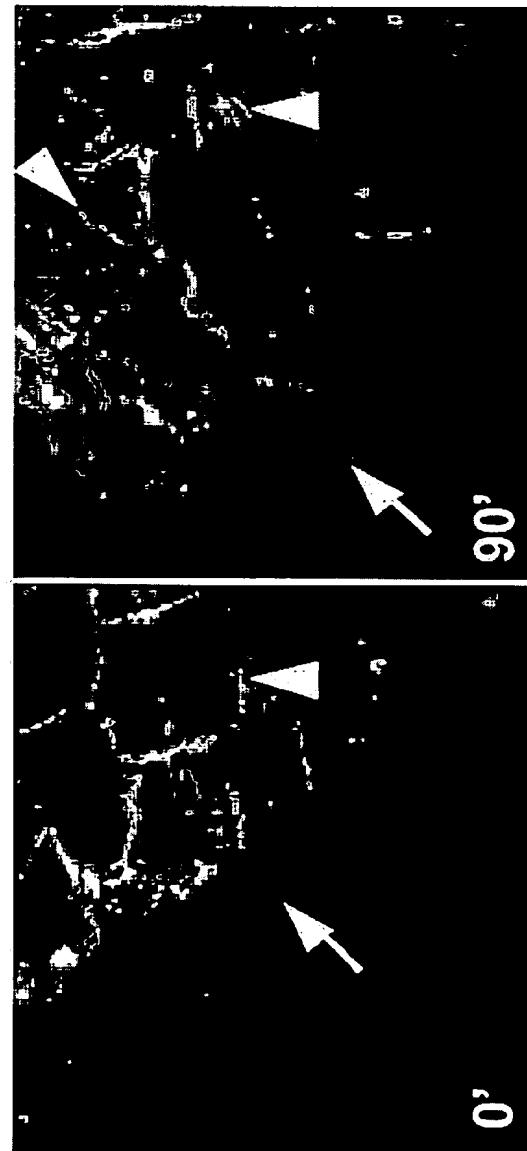


FIG. 17B

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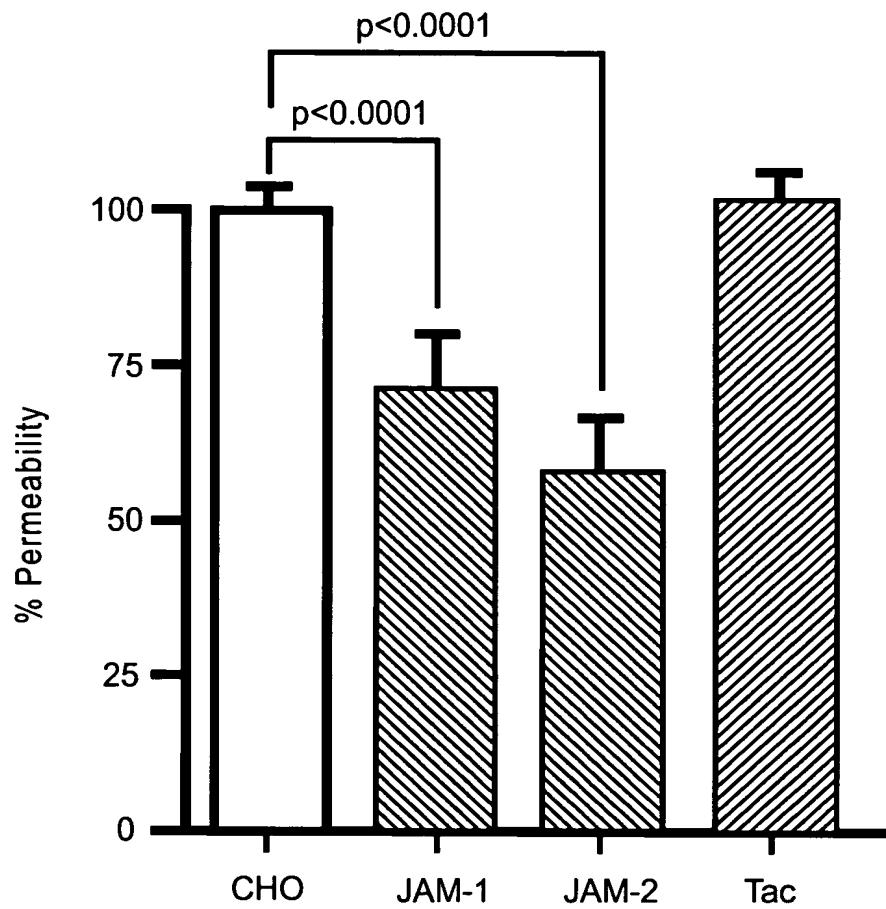


FIG. 18



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BY	CLASS
DRAFTSMAN	SUBCLASS

→ Apical
Basal

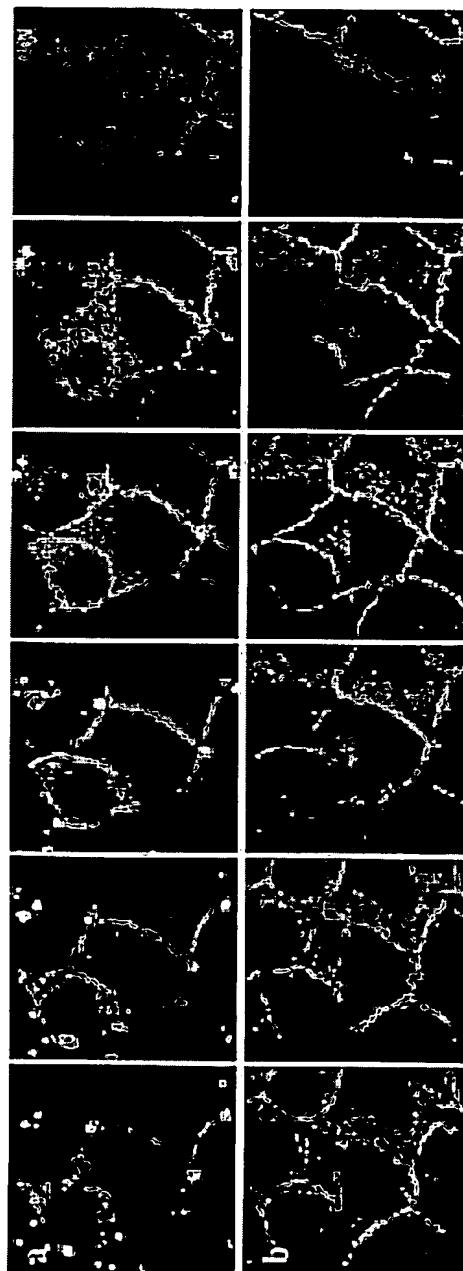


FIG. 19A

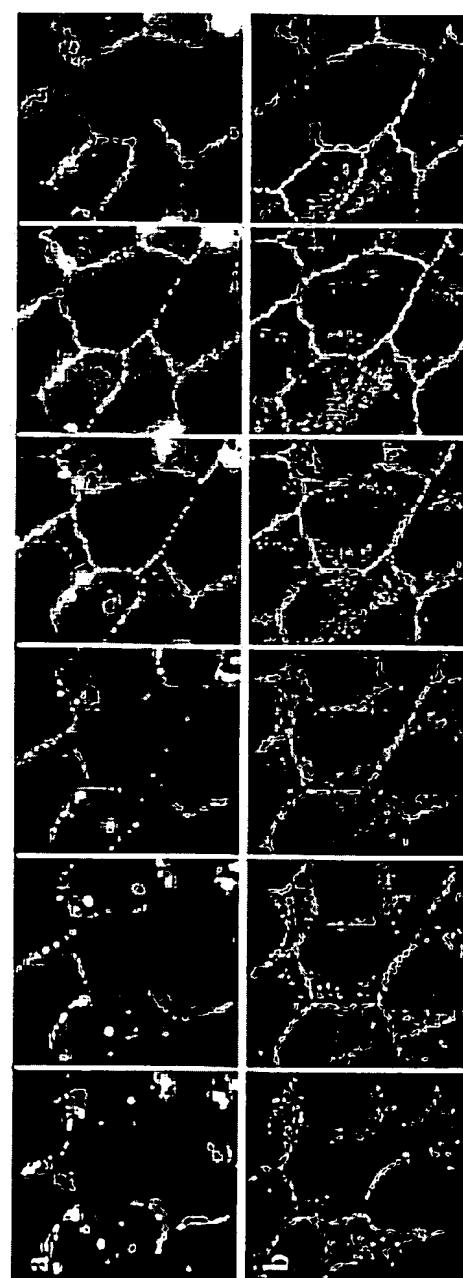


FIG. 19B

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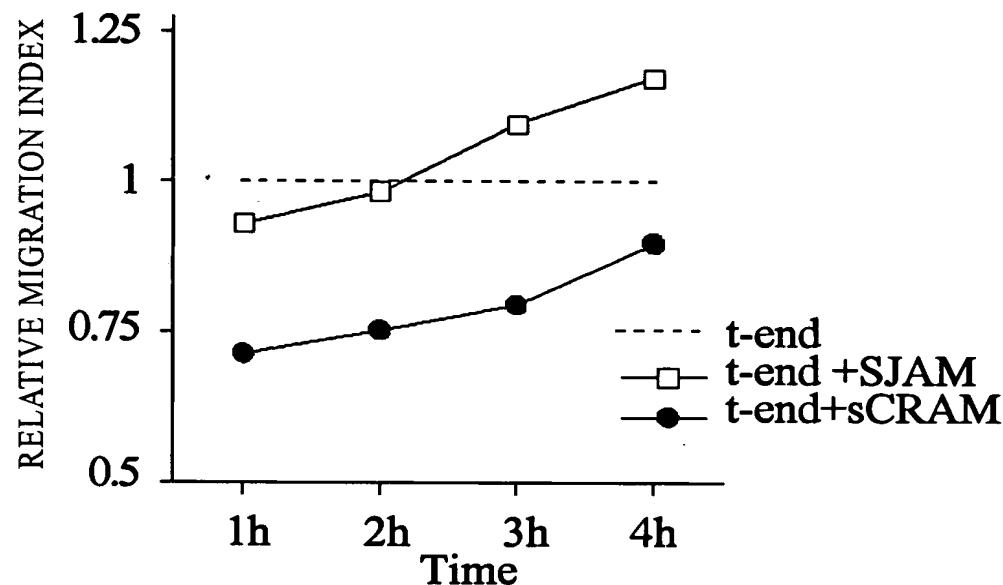
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FIG. 20A



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FIG. 20B

